

## WEST Search History

DATE: Wednesday, October 09, 2002

<u>Set Name</u> side by side	<u>Query</u>	<u>Hit Count</u>	<u>Set Name</u> result set
<i>DB=USPT,PGPB; PLUR=YES; OP=ADJ</i>			
L9	nitrate responsive	2	L9
L8	11 and lateral root	35	L8
L7	15 and lateral root	9	L7
L6	15 and lateral root	9	L6
L5	14 and maize	78	L5
L4	L3 and transcription factor	88	L4
L3	L1 and transgenic	711	L3
L2	L1 and anr1	1	L2
L1	nitrate and root	5385	L1

END OF SEARCH HISTORY

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NEWS 5 Apr 19 US Patent Applications available in IFICDB, IFIPAT, and IFIUDB  
NEWS 6 Apr 22 Records from IP.com available in CAPLUS, HCAPLUS, and ZCAPLUS  
NEWS 7 Apr 22 BIOSIS Gene Names now available in TOXCENTER  
NEWS 8 Apr 22 Federal Research in Progress (FEDRIP) now available  
NEWS 9 Jun 03 New e-mail delivery for search results now available  
NEWS 10 Jun 10 MEDLINE Reload  
NEWS 11 Jun 10 PCTFULL has been reloaded  
NEWS 12 Jul 02 FOREGE no longer contains STANDARDS file segment  
NEWS 13 Jul 22 USAN to be reloaded July 28, 2002;  
saved answer sets no longer valid  
NEWS 14 Jul 29 Enhanced polymer searching in REGISTRY  
NEWS 15 Jul 30 NETFIRST to be removed from STN  
NEWS 16 Aug 08 CANCERLIT reload  
NEWS 17 Aug 08 PHARMAMarketLetter(PHARMAML) - new on STN  
NEWS 18 Aug 08 NTIS has been reloaded and enhanced  
NEWS 19 Aug 19 Aquatic Toxicity Information Retrieval (AQUIRE)  
now available on STN  
NEWS 20 Aug 19 IFIPAT, IFICDB, and IFIUDB have been reloaded  
NEWS 21 Aug 19 The MEDLINE file segment of TOXCENTER has been reloaded  
NEWS 22 Aug 26 Sequence searching in REGISTRY enhanced  
NEWS 23 Sep 03 JAPIO has been reloaded and enhanced  
NEWS 24 Sep 16 Experimental properties added to the REGISTRY file  
NEWS 25 Sep 16 Indexing added to some pre-1967 records in CA/CAPLUS  
NEWS 26 Sep 16 CA Section Thesaurus available in CAPLUS and CA  
NEWS 27 Oct 01 CASREACT Enriched with Reactions from 1907 to 1985  
  
NEWS EXPRESS February 1 CURRENT WINDOWS VERSION IS V6.0d,  
CURRENT MACINTOSH VERSION IS V6.0a(ENG) AND V6.0Ja(JP),  
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FILE 'HOME' ENTERED AT 11:28:43 ON 09 OCT 2002

=> file agricola caplus biosis

COST IN U.S. DOLLARS

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SESSION

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0.21

0.21

FILE 'AGRICOLA' ENTERED AT 11:28:53 ON 09 OCT 2002

FILE 'CAPLUS' ENTERED AT 11:28:53 ON 09 OCT 2002

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FILE 'BIOSIS' ENTERED AT 11:28:53 ON 09 OCT 2002

COPYRIGHT (C) 2002 BIOLOGICAL ABSTRACTS INC. (R)

=> s nitrate responsive

L1 44 NITRATE RESPONSIVE

=> s l1 and plant?

L2 17 L1 AND PLANT?

=> dup rem l2

PROCESSING COMPLETED FOR L2

L3 8 DUP REM L2 (9 DUPLICATES REMOVED)

=> d 1-8 ti

L3 ANSWER 1 OF 8 CAPLUS COPYRIGHT 2002 ACS

TI Discovery of gene for maize **nitrate-responsive** root transcription factor sequence homologs for control of root development in transgenic **plants**

L3 ANSWER 2 OF 8 AGRICOLA

DUPPLICATE 1

TI Nitrate-induced genes in tomato roots. Array analysis reveals novel genes that may play a role in nitrogen nutrition. [Erratum: Nov 2001, v. 127 (3), p. 1323.]

L3 ANSWER 3 OF 8 AGRICOLA

DUPPLICATE 2

TI Identification of light- and **nitrate-responsive** regions of the nitrate reductase promoter from birch.

L3 ANSWER 4 OF 8 CAPLUS COPYRIGHT 2002 ACS

DUPPLICATE 3

TI Functional analysis of a nitrite reductase promoter from birch in transgenic tobacco

L3 ANSWER 5 OF 8 CAPLUS COPYRIGHT 2002 ACS

TI Cloning of cDNA for cytokinin-inducible protein 1 from maize

L3 ANSWER 6 OF 8 AGRICOLA

DUPPLICATE 4

TI Differential expression of genes for response regulators in response to cytokinins and nitrate in *Arabidopsis thaliana*.

L3 ANSWER 7 OF 8 AGRICOLA

DUPPLICATE 5

TI A response-regulator homologue possibly involved in nitrogen signal transduction mediated by cytokinin in maize.

L3 ANSWER 8 OF 8 CAPLUS COPYRIGHT 2002 ACS

TI 5' Proximal regions of *Arabidopsis* nitrate reductase genes direct

nitrate-induced transcription in transgenic tobacco

=> d ab

L3 ANSWER 1 OF 8 CAPLUS COPYRIGHT 2002 ACS

AB The invention provides isolated maize **nitrate-responsive** root transcription factor nucleic acids, identified by sequence homol., and their encoded proteins. The present invention provides methods and compns. relating to altering root transcriptional factor levels in transgenic **plants**. The invention further provides recombinant expression cassettes, host cells, and transgenic **plants**.

=> d pi

L3 ANSWER 1 OF 8 CAPLUS COPYRIGHT 2002 ACS

PATENT NO. KIND DATE APPLICATION NO. DATE

PI	WO 2002029069	A2	20020411	WO 2001-US30814	20011003	
	W:	AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM				
	RW:	GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR				
AU	2002011353	A5	20020415	AU 2002-11353	20011003	
US	2002124284	A1	20020905	US 2001-970624	20011004	

=> d 2 ab

L3 ANSWER 2 OF 8 AGRICOLA

DUPLICATE 1

AB A subtractive tomato (*Lycopersicon esculentum*) root cDNA library enriched in genes up-regulated by changes in **plant** mineral status was screened with labeled mRNA from roots of both nitrate-induced and mineral nutrient-deficient (-nitrogen [N], -phosphorus, -potassium [K], -sulfur, -magnesium, -calcium, -iron, -zinc, and -copper) tomato **plants**. A subset of cDNAs was selected from this library based on mineral nutrient-related changes in expression. Additional cDNAs were selected from a second mineral-deficient tomato root library based on sequence homology to known genes. These selection processes yielded a set of 1,280 mineral nutrition-related cDNAs that were arrayed on nylon membranes for further analysis. These high-density arrays were hybridized with mRNA from tomato **plants** exposed to nitrate at different time points after N was withheld for 48 h, for **plants** that were grown on nitrate/ammonium for 5 weeks prior to the withholding of N. One hundred-fifteen genes were found to be up-regulated by nitrate resupply. Among these genes were several previously identified as **nitrate responsive**, including nitrate transporters, nitrate and nitrite reductase, and metabolic enzymes such as transaldolase, transketolase, malate dehydrogenase, asparagine synthetase, and histidine decarboxylase. We also identified 14 novel nitrate-inducible genes, including: (a) water channels, (b) root phosphate and K<sup>+</sup> transporters, (c) genes potentially involved in transcriptional regulation, (d) stress response genes, and (e) ribosomal protein genes. In addition, both families of nitrate transporters were also found to be inducible by phosphate, K, and iron deficiencies. The identification of these novel nitrate-inducible genes is providing avenues of research that will yield new insights into the molecular basis of **plant** N nutrition, as well as possible networking between the regulation of N, phosphorus, and K nutrition.

=> d 2 so

L3 ANSWER 2 OF 8 AGRICOLA DUPLICATE 1  
SO Plant physiology, Sept 2001. Vol. 127, No. 1. p. 345-359  
Publisher: Rockville, MD : American Society of Plant Physiologists, 1926-  
CODEN: PLPHAY; ISSN: 0032-0889

=> d 6 ab

L3 ANSWER 6 OF 8 AGRICOLA DUPLICATE 4  
AB In *Arabidopsis thaliana*, a number of response regulators are presumably involved in His-Asp phosphorelay signal transduction in response to environmental stimuli, such as phytohormones. Previously, it was shown that expression of a certain set of genes for response regulators are cytokinin- and **nitrate-responsive** in their mRNA accumulation, under certain growth conditions [Taniguchi et al. (1998) FEBS Lett. 429: 259, Brandstatter and Kieber (1998) *Plant Cell* 10: 1009]. To answer the critical question of whether or not other response regulator genes, so far identified in *Arabidopsis thaliana*, are also cytokinin-inducible, here an extended comparative examination was carried out. It was demonstrated that not all of response regulator genes are necessarily cytokinin-responsive in their transcription. Rather, the members of a certain subfamily (type-A) are cytokinin-responsive, but those belonging to the other (type-B) are not. The presumed nitrate-responsiveness was also assessed for the same set of response regulators, and the analogous view was supported. These results suggest that the two subtypes of response regulators differ from each other, as judged from not only their structural designs, but also the expression profiles of their transcripts in response to **plant** stimuli.

=> d 7 ab

L3 ANSWER 7 OF 8 AGRICOLA DUPLICATE 5  
AB A cDNA clone, pZmCip1, encoding a maize (*Zea mays*) cytokinin-inducible protein 1 was isolated utilizing the differential display technique, and studied using the expression of ZmCip1 in nitrogen-starved maize **plants**. The cloned cDNA contained an open reading frame consisting of 157 amino acids with a predicted molecular mass of 16.7 kDa, which possesses similarity with the response-regulators of bacterial two-component signalling systems. In detached leaves, accumulation of ZmCip1 transcript by t-zeatin was dose-dependent in a range of 10(-9) M to 10(-7) M, and occurred within 30 min after treatment. The effect of t-zeatin was replaceable by isopentenyl-adenosine or isopentenyl-adenosine-5'-monophosphate. Pretreatment of detached leaves with cycloheximide did not inhibit the accumulation of the transcript. In whole **plants**, ZmCip1 transcript was transiently accumulated exclusively in leaves by supply of nitrate or ammonium ions to the roots, whereas the transcript was not accumulated in detached leaves by supply of the nitrogen nutrients. Both the cytokinin- and **nitrate-responsive** accumulations of ZmCip1 transcript were accompanied by an increase in the immunotitratable protein. Isopentenyladenosine and/or its phosphorylated form(s) accumulated in roots 2 h after supply of nitrate to **plants**. These results, taken together, suggest that ZmCip1 is a primary response gene to cytokinins, and that it involves, at least in part, the nitrogen-signal transduction mediated by cytokinin in maize.

=> s nitrate and root?

L4 15273 NITRATE AND ROOT?

=> s 14 and lateral root?

L5 193 L4 AND LATERAL ROOT?

=> s 15 and (gene or cdna or coding region)

L6 31 L5 AND (GENE OR CDNA OR CODING REGION)

=> dup rem 16

PROCESSING COMPLETED FOR L6

L7 15 DUP REM L6 (16 DUPLICATES REMOVED)

=> d 1-15 ti

L7 ANSWER 1 OF 15 CAPLUS COPYRIGHT 2002 ACS DUPLICATE 1

TI The novel symbiotic phenotype of enhanced-nodulating mutant of *Lotus japonicus*: astray mutant is an early nodulating mutant with wider nodulation zone

L7 ANSWER 2 OF 15 CAPLUS COPYRIGHT 2002 ACS DUPLICATE 2

TI The *Arabidopsis* dual-affinity **nitrate transporter gene** AtNRT1.1 (CHL1) is regulated by auxin in both shoots and roots

L7 ANSWER 3 OF 15 CAPLUS COPYRIGHT 2002 ACS DUPLICATE 3

TI AUX1 promotes **lateral root** formation by facilitating indole-3-acetic acid distribution between sink and source tissues in the *arabidopsis* seedling

L7 ANSWER 4 OF 15 AGRICOLA DUPLICATE 4

TI The *Arabidopsis* dual-affinity **nitrate transporter gene** AtNRT1.1 (CHL1) is activated and functions in nascent organ development during vegetative and reproductive growth.

L7 ANSWER 5 OF 15 CAPLUS COPYRIGHT 2002 ACS

TI Soil and plant specific effects on bacterial community composition in the rhizosphere

L7 ANSWER 6 OF 15 BIOSIS COPYRIGHT 2002 BIOLOGICAL ABSTRACTS INC.

TI ABA plays a central role in mediating the regulatory effects of **nitrate** on **root** branching in *Arabidopsis*.

L7 ANSWER 7 OF 15 CAPLUS COPYRIGHT 2002 ACS DUPLICATE 5

TI Control of plant development by limiting factors: a nutritional perspective

L7 ANSWER 8 OF 15 CAPLUS COPYRIGHT 2002 ACS DUPLICATE 6

TI Impact of *Agrobacterium tumefaciens*-induced stem tumors on NO<sub>3</sub><sup>-</sup> uptake in *Ricinus communis*

L7 ANSWER 9 OF 15 AGRICOLA DUPLICATE 7

TI Regulation of *Arabidopsis* **root** development by **nitrate** availability.

L7 ANSWER 10 OF 15 CAPLUS COPYRIGHT 2002 ACS

TI **Nitrate** acts as a signal to control **gene** expression, metabolism and biomass allocation

L7 ANSWER 11 OF 15 AGRICOLA DUPLICATE 8

TI Evidence for **nitrate** reductase expression during initiation of **lateral roots** by NAA in chicory.

L7 ANSWER 12 OF 15 AGRICOLA DUPLICATE 9

TI Expression studies of Nrt2:1Np, a putative high-affinity **nitrate** transporter: evidence for its role in **nitrate** uptake.

L7 ANSWER 13 OF 15 AGRICOLA DUPLICATE 10

TI An *Arabidopsis* MADS box **gene** that controls nutrient-induced

changes in **root** architecture.

L7 ANSWER 14 OF 15 CAPLUS COPYRIGHT 2002 ACS  
TI Molecular and genetic insights into shoot control of nodulation in soybean

L7 ANSWER 15 OF 15 CAPLUS COPYRIGHT 2002 ACS DUPLICATE 11  
TI Studies on the **root** control of non-nodulation and plant growth  
of non-nodulating mutants and a supernodulating mutant of soybean (*Glycine*  
*max* (L.) Merr.)

=> d 3 ab

L7 ANSWER 3 OF 15 CAPLUS COPYRIGHT 2002 ACS DUPLICATE 3  
AB *Arabidopsis* **root** architecture is regulated by shoot-derived  
signals such as **nitrate** and auxin. We report that mutations in  
the putative auxin influx carrier AUX1 modify **root** architecture  
as a result of the disruption in hormone transport between indole-3-acetic  
acid (IAA) source and sink tissues. Gas chromatog.-selected reaction  
monitoring-mass spectrometry measurements revealed that the aux1 mutant  
exhibited altered IAA distribution in young leaf and **root**  
tissues, the major IAA source and sink organs, resp., in the developing  
seedling. Expression studies using the auxin-inducible reporter  
IAA2::uidA revealed that AUX1 facilitates IAA loading into the leaf  
vascular transport system. AUX1 also facilitates IAA unloading in the  
primary **root** apex and developing **lateral root**  
primordium. Exogenous application of the synthetic auxin 1-naphthylacetic  
acid is able to rescue the aux1 **lateral root**  
phenotype, implying that **root** auxin levels are suboptimal for  
**lateral root** primordium initiation in the mutant.

=> d 4 ab

L7 ANSWER 4 OF 15 AGRICOLA DUPLICATE 4  
AB The AtNRT1.1 (CHL1) transporter provides a primary mechanism for  
**nitrate** uptake in *Arabidopsis* and is expected to localize to the  
epidermis and cortex of the mature **root**, where the bulk of  
**nitrate** uptake occurs. Using fusions to GFP/GUS marker genes, we  
found CHL1 expression concentrated in the tips of primary and  
**lateral roots**, with very low signals in the epidermis  
and cortex. A time-course study showed that CHL1 is activated in the  
primary **root** tip early in seedling development and at the  
earliest stages of **lateral root** formation. Strong CHL1  
expression also was found in shoots, concentrated in young leaves and  
developing flower buds but not in the shoot meristem. These expression  
patterns were confirmed by immunolocalization and led us to examine CHL1  
function specifically in the growth of developing organs. chl1 mutants  
showed a reduction in the growth of nascent **roots**, stems,  
leaves, and flower buds. The growth of nascent primary **roots** was  
inhibited in the mutants even in the absence of added **nitrate**,  
whereas elongation of **lateral root** primordia was  
inhibited specifically at low **nitrate** and acidic pH.  
Interestingly, chl1 mutants also displayed a late-flowering phenotype.  
These results indicate that CHL1 is activated and functions in the growth  
of nascent organs in both shoots and **roots** during vegetative and  
reproductive growth.

=> d 4 so

L7 ANSWER 4 OF 15 AGRICOLA DUPLICATE 4  
SO The Plant cell, Aug 2001. Vol. 13, No. 8. p. 1761-1777  
Publisher: [Rockville, MD : American Society of Plant Physiologists,

c1989-  
CODEN: PLCEEW; ISSN: 1040-4651

=> d 6 ab

L7 ANSWER 6 OF 15 BIOSIS COPYRIGHT 2002 BIOLOGICAL ABSTRACTS INC.  
AB The formation of **lateral roots** (LR) is a major post-embryonic developmental event in plants. In *Arabidopsis thaliana*, LR development is inhibited by high concentrations of NO<sub>3</sub><sup>-</sup>. Here we present strong evidence that ABA plays an important role in mediating the effects of NO<sub>3</sub><sup>-</sup> on LR formation. Firstly, the inhibitory effect of NO<sub>3</sub><sup>-</sup> is significantly reduced in three ABA insensitive mutants, *abi4-1*, *abi4-2* and *abi5-1*, but not in *abi1-1* and *abi3-1*. Secondly, inhibition by NO<sub>3</sub><sup>-</sup> is significantly reduced, but not completely abolished, in four ABA synthesis mutants, *aba1-1*, *aba2-3*, *aba2-4* and *aba3-2*. These results indicate that there are two regulatory pathways mediating the inhibitory effects of NO<sub>3</sub><sup>-</sup> in *A. thaliana* **roots**. One pathway is ABA-dependent and involves *ABI4* and *ABI5*, whereas the second pathway is ABA-independent. In addition, ABA also plays a role in mediating the stimulation of LR elongation by local NO<sub>3</sub><sup>-</sup> applications.

=> d 6 so

L7 ANSWER 6 OF 15 BIOSIS COPYRIGHT 2002 BIOLOGICAL ABSTRACTS INC.  
SO Plant Journal, (December, 2001) Vol. 28, No. 6, pp. 655-662.  
<http://www.blackwell-science.com/cgilib/jnlpage.bin?Journal=TPJ&File=TPJ&Page=aims.print>.  
ISSN: 0960-7412.

=> d 13 ab

L7 ANSWER 13 OF 15 AGRICOLA DUPLICATE 10  
AB The development of plant **root** systems is sensitive to the availability and distribution of nutrients within the soil. For example, **lateral roots** proliferate preferentially within **nitrate** (NO<sub>3</sub><sup>-</sup>)-rich soil patches. A NO<sub>3</sub><sup>-</sup>-inducible *Arabidopsis* **gene** (ANR1), was identified that encodes a member of the MADS box family of transcription factors. Transgenic plants in which ANR1 was repressed had an altered sensitivity to NO<sub>3</sub><sup>-</sup> and no longer responded to NO<sub>3</sub><sup>-</sup>-rich zones by **lateral root** proliferation, indicating that ANR1 is a key determinant of developmental plasticity in *Arabidopsis* **roots**.

=> d 13 so

L7 ANSWER 13 OF 15 AGRICOLA DUPLICATE 10  
SO Science, Jan 16, 1998. Vol. 279, No. 5349. p. 407-409  
Publisher: Washington, D.C. : American Association for the Advancement of Science.  
CODEN: SCIEAS; ISSN: 0036-8075

=> dis his

(FILE 'HOME' ENTERED AT 11:28:43 ON 09 OCT 2002)

FILE 'AGRICOLA, CAPLUS, BIOSIS' ENTERED AT 11:28:53 ON 09 OCT 2002  
L1 44 S NITRATE RESPONSIVE  
L2 17 S L1 AND PLANT?  
L3 8 DUP REM L2 (9 DUPLICATES REMOVED)

L4 15273 S NITRATE AND ROOT?  
 L5 193 S L4 AND LATERAL ROOT?  
 L6 31 S L5 AND (GENE OR CDNA OR CODING REGION)  
 L7 15 DUP REM L6 (16 DUPLICATES REMOVED)

=> s nitrate and transcription factor and root?  
 L8 4 NITRATE AND TRANSCRIPTION FACTOR AND ROOT?

=> dup rem 18  
 PROCESSING COMPLETED FOR L8  
 L9 3 DUP REM L8 (1 DUPLICATE REMOVED)

=> d 1-3 ti

L9 ANSWER 1 OF 3 CAPLUS COPYRIGHT 2002 ACS  
 TI Discovery of gene for maize **nitrate-responsive root transcription factor** sequence homologs for control of root development in transgenic plants

L9 ANSWER 2 OF 3 CAPLUS COPYRIGHT 2002 ACS  
 TI Sequence-determined DNA fragments and corresponding encoded polypeptides from corn and Arabidopsis

L9 ANSWER 3 OF 3 CAPLUS COPYRIGHT 2002 ACS DUPLICATE 1  
 TI Azorhizobium caulinodans nitrogen fixation (nif/fix) gene regulation: mutagenesis of the nifA -24/-12 promoter element, characterization of a ntrA(rpoN) gene, and derivation of a model

=> d 2 ab

L9 ANSWER 2 OF 3 CAPLUS COPYRIGHT 2002 ACS  
 AB The present invention provides DNA mols. that constitute fragments of the genome and cDNAs from Zea mays mays (HYBRID SEED #35A19) and Arabidopsis thaliana (ecotype Wassilewski), and polypeptides encoded thereby. The DNA mols. are useful for specifying a gene product in cells, either as a promoter or as a protein coding sequence or as an UTR or as a 3' termination sequence, and are also useful in controlling the behavior of a gene in the chromosome, in controlling the expression of a gene or as tools for genetic mapping, recognizing or isolating identical or related DNA fragments, or identification of a particular individual organism, or for clustering of a group of organisms with a common trait. Arabidopsis DNA is used in the present expt., but the procedure is a general one. Protocols are provided for Southern hybridizations and transformation of carrot cells. [This abstr. record is one of 15 records supplemental to CA13316218528Q necessitated by the large no. of index entries required to fully index the document and publication system constraints.].

=> d 2 pi

L9 ANSWER 2 OF 3 CAPLUS COPYRIGHT 2002 ACS  
 PATENT NO. KIND DATE APPLICATION NO. DATE  
 -----  
 PI EP 1033405 A2 20000906 EP 2000-301439 20000225  
 R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,  
 IE, SI, LT, LV, FI, RO  
 CA 2300692 AA 20000825 CA 2000-2300692 20000225  
 CA 2302828 AA 20001006 CA 2000-2302828 20000406  
 EP 1055728 A2 20001129 EP 2000-303770 20000504  
 R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,  
 IE, SI, LT, LV, FI, RO  
 EP 1054060 A2 20001122 EP 2000-304161 20000517  
 R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,

IE, SI, LT, LV, FI, RO

=> s anrl

L10 25 ANR1

=> s l10 and transgenic

L11 3 L10 AND TRANSGENIC

=> dup rem l11

PROCESSING COMPLETED FOR L11

L12 1 DUP REM L11 (2 DUPLICATES REMOVED)

=> d ti

L12 ANSWER 1 OF 1 AGRICOLA

DUPLICATE 1

TI An Arabidopsis MADS box gene that controls nutrient-induced changes in root architecture.

=> d so

L12 ANSWER 1 OF 1 AGRICOLA

DUPLICATE 1

SO Science, Jan 16, 1998. Vol. 279, No. 5349. p. 407-409

Publisher: Washington, D.C. : American Association for the Advancement of Science.

CODEN: SCIEAS; ISSN: 0036-8075

=> dup rem l10

PROCESSING COMPLETED FOR L10

L13 15 DUP REM L10 (10 DUPLICATES REMOVED)

=> s l10 and plant?

L14 14 L10 AND PLANT?

=> s l13 and plant?

L15 5 L13 AND PLANT?

=> d 1-5 ti

L15 ANSWER 1 OF 5 AGRICOLA

TI Genomic analysis of a nutrient response in arabidopsis reveals diverse expression patterns and novel metabolic and potential regulatory genes induced by nitrate.

L15 ANSWER 2 OF 5 AGRICOLA

TI Regulation of Arabidopsis root development by nitrate availability.

L15 ANSWER 3 OF 5 AGRICOLA

TI An Arabidopsis MADS box gene that controls nutrient-induced changes in root architecture.

L15 ANSWER 4 OF 5 BIOSIS COPYRIGHT 2002 BIOLOGICAL ABSTRACTS INC.

TI Environmental factors controlling flower opening and closing in a Portulaca hybrid.

L15 ANSWER 5 OF 5 BIOSIS COPYRIGHT 2002 BIOLOGICAL ABSTRACTS INC.

TI Functional analysis of a novel nitrogen-inducible MADS-box gene expressed in Arabidopsis roots.

=> d 5 so

L15 ANSWER 5 OF 5 BIOSIS COPYRIGHT 2002 BIOLOGICAL ABSTRACTS INC.  
SO Plant Physiology (Rockville), (1997) Vol. 114, No. 3 SUPPL., pp. 72.  
Meeting Info.: PLANT BIOLOGY '97: 1997 Annual Meetings of the American  
Society of Plant Physiologists and the Canadian Society of Plant  
Physiologists, Japanese Society of Plant Physiologists and the Australian  
Society of Plant Physiologists Vancouver, British Columbia, Canada August  
2-6, 1997  
ISSN: 0032-0889.

=> d 5 ab

L15 ANSWER 5 OF 5 BIOSIS COPYRIGHT 2002 BIOLOGICAL ABSTRACTS INC.

=> d 5 au

L15 ANSWER 5 OF 5 BIOSIS COPYRIGHT 2002 BIOLOGICAL ABSTRACTS INC.  
AU Zhang, Hanma; Forde, Brian G.

=> d 3 ab

L15 ANSWER 3 OF 5 AGRICOLA

AB The development of **plant** root systems is sensitive to the availability and distribution of nutrients within the soil. For example, lateral roots proliferate preferentially within nitrate (NO<sub>3</sub><sup>-</sup>)-rich soil patches. A NO<sub>3</sub><sup>-</sup>-inducible *Arabidopsis* gene (**ANR1**), was identified that encodes a member of the MADS box family of transcription factors. Transgenic **plants** in which **ANR1** was repressed had an altered sensitivity to NO<sub>3</sub><sup>-</sup> and no longer responded to NO<sub>3</sub><sup>-</sup>-rich zones by lateral root proliferation, indicating that **ANR1** is a key determinant of developmental plasticity in *Arabidopsis* roots.

=> d 3 so

L15 ANSWER 3 OF 5 AGRICOLA

SO Science, Jan 16, 1998. Vol. 279, No. 5349. p. 407-409  
Publisher: Washington, D.C. : American Association for the Advancement of  
Science.  
CODEN: SCIEAS; ISSN: 0036-8075

=> d ab

L15 ANSWER 1 OF 5 AGRICOLA

AB Microarray and RNA gel blot analyses were performed to identify *Arabidopsis* genes that responded to nitrate at both low (250 micromolar) and high (5 to 10 mM) nitrate concentrations. Genes involved directly or indirectly with nitrite reduction were the most highly induced by nitrate. Most of the known nitrate-regulated genes (including those encoding nitrate reductase, the nitrate transporter NRT1, and glutamate synthase) appeared in the 40 most strongly nitrate-induced genes/clones on at least one of the microarrays of the 5524 genes/clones investigated. Novel nitrate-induced genes were also found, including those encoding (1) possible regulatory proteins, including an MYB transcription factor, a calcium antiporter, and putative protein kinases; (2) metabolic enzymes, including transaldolase and transketolase of the nonoxidative pentose pathway, malate dehydrogenase, asparagine synthetase, and histidine decarboxylase; and (3) proteins with unknown functions, including nonsymbiotic hemoglobin, a senescence-associated protein, and two methyl-transferases. The primary pattern of induction observed for many of these genes was a transient increase in mRNA at low nitrate concentrations

and a sustained increase when treated with high nitrate concentrations. Other patterns of induction observed included transient inductions after both low and high nitrate treatments and sustained or increasing amounts of mRNA after either treatment. Two genes, *AMT1;1* encoding an ammonium transporter and **ANR1** encoding a MADS-box factor, were repressed by nitrate. These findings indicate that nitrate induces not just one but many diverse responses at the mRNA level in *Arabidopsis*.

=> d so

L15 ANSWER 1 OF 5 AGRICOLA

SO The Plant cell, Aug 2000. Vol. 12, No. 8. p. 1491-1509

Publisher: [Rockville, MD : American Society of Plant Physiologists, c1989-

CODEN: PLCEEW; ISSN: 1040-4651

=> d au

L15 ANSWER 1 OF 5 AGRICOLA

AU Wang, R.; Guegler, K.; LaBrie, S.T.; Crawford, N.M.

=> s maize and nitrate and root?

L16 1016 MAIZE AND NITRATE AND ROOT?

=> s l16 and transcription factor

L17 1 L16 AND TRANSCRIPTION FACTOR

=> d ti

L17 ANSWER 1 OF 1 CAPLUS COPYRIGHT 2002 ACS

TI Discovery of gene for maize nitrate-responsive root transcription factor sequence homologs for control of root development in transgenic plants

=> d pi

L17 ANSWER 1 OF 1 CAPLUS COPYRIGHT 2002 ACS

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
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PI	WO 2002029069	A2	20020411	WO 2001-US30814 20011003
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W:	AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM
RW:	GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR

AU 2002011353	A5	20020415	AU 2002-11353	20011003
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US 2002124284	A1	20020905	US 2001-970624	20011004
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=> s ((bruce w?) or (bruce, w?))/au

L18 838 ((BRUCE W?) OR (BRUCE, W?))/AU

=> s l18 and nitrate

L19 13 L18 AND NITRATE

=> dup rem l19

PROCESSING COMPLETED FOR L19

L20 7 DUP REM L19 (6 DUPLICATES REMOVED)

=> d 1-7 ti

L20 ANSWER 1 OF 7 CAPLUS COPYRIGHT 2002 ACS  
TI Discovery of gene for maize **nitrate**-responsive root  
transcription factor sequence homologs for control of root development in  
transgenic plants

L20 ANSWER 2 OF 7 CAPLUS COPYRIGHT 2002 ACS DUPLICATE 1  
TI Analysis and formation of nitrosamines in the human intestine

L20 ANSWER 3 OF 7 CAPLUS COPYRIGHT 2002 ACS DUPLICATE 2  
TI Absence of volatile nitrosamines in human feces

L20 ANSWER 4 OF 7 CAPLUS COPYRIGHT 2002 ACS DUPLICATE 3  
TI Reevaluation of **nitrate** and nitrite levels in the human  
intestine

L20 ANSWER 5 OF 7 AGRICOLA DUPLICATE 4  
TI Analysis of **nitrate**, nitrite, and nitrosamines in human feces.

L20 ANSWER 6 OF 7 CAPLUS COPYRIGHT 2002 ACS DUPLICATE 5  
TI Nitrite and **nitrate** are formed by endogenous synthesis in the  
human intestine

L20 ANSWER 7 OF 7 CAPLUS COPYRIGHT 2002 ACS  
TI (Benzylideneamino)guanidines